



**Figure S14. Normalized distribution of complexes in the DOCKGROUND benchmark set 3 according to TM performance,  $P_{TM}$  (Eq. 1).** The data is obtained by the basic TM protocol with OR-queries, and is normalized to the total number of complexes, for which residues were predicted (column 3 in Table 3). Dark and light bars show distributions for all retrieved residues and for the top 10 residues, submitted to docking, respectively.